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SEQUENCE LISTING

<110> Istituto Superiore di Sanità <120> Novel Tat Complexes, And Vaccines Comprising Them <130> WPP89320 <150> UK 0405480.5 <151> 2004-03-11 <160> 2 <170> PatentIn version 3.3 <210> 1 <211> 102 <212> PRT <213> Human immunodeficiency virus type 1 <400> 1 Met Glu Pro Val Asp Pro Arg Leu Glu Pro Trp Lys His Pro Gly Ser Gln Pro Lys Thr Ala Cys Thr Asn Cys Tyr Cys Lys Lys Cys Cys Phe 25 His Cys Gln Val Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly 40 Arg Lys Lys Arg Arg Gln Arg Arg Pro Pro Gln Gly Ser Gln Thr 55 His Gln Val Ser Leu Ser Lys Gln Pro Thr Ser Gln Ser Arg Gly Asp 65 Pro Thr Gly Pro Lys Glu Gln Lys Lys Lys Val Glu Arg Glu Thr Glu Thr Asp Pro Val His Gln 100 <210> 2 <211> 853 <212> PRT <213> Human immunodeficiency virus type 1 <220> <221> misc_feature <222> (23)..(23) <223> Xaa can be any naturally occurring amino acid <220> <221> misc_feature <222> (84)..(86) <223> Xaa can be any naturally occurring amino acid

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Gln Ser Leu Lys Pro Cys Val Lys Leu Thr Pro Leu C 115 120 1	Cys Val Thr Leu 125
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165 170 175

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Gly Phe Ala Ile Leu Lys Cys Lys Asp Lys Lys Phe Asn Gly Lys Gly 225 230 235

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Asp Pro Glu Ile Val Thr His Ser Phe Asn Cys Gly Glu Phe Phe 370 375 380

Tyr Cys Asn Ser Thr Gln Leu Phe Asn Ser Thr Trp Asn Val Thr Glu 385 390 395 400

Glu Ser Asn Asn Thr Val Glu Asn Asn Thr Ile Thr Leu Pro Cys Arg 405 410 415

Ile Lys Gln Ile Ile Asn Met Trp Gln Xaa Val Gly Arg Ala Met Tyr 420 425 430

Ala Pro Pro Ile Arg Gly Gln Ile Arg Cys Ser Ser Asn Ile Thr Gly

4

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435 440 445

Leu Leu Thr Arg Asp Gly Gly Pro-Glu Asp Asn Lys Thr Glu Val 450 455 460

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Phe Arg Pro Gly Gly Gly Asp Met Arg Asp Asn Trp Arg Ser Glu Leu 465 470 475

Tyr Lys Tyr Lys Val Val Lys Ile Glu Pro Leu Gly Val Ala Pro Thr 485 490 495

Lys Ala Lys Arg Arg Val Val Gln Arg Glu Lys Arg Ala Val Gly Ile 500 505 510

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Ala Ala Ser Met Thr Leu Thr Val Gln Ala Arg Leu Leu Ser Gly 530 535 540

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His Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg
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Val Leu Ala Val Glu Arg Tyr Leu Arg Asp Gln Gln Leu Leu Gly Ile 580 585 590

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Ala Ser Trp Ser Asn Lys Ser Leu Asn Lys Ile Trp Asp Asn Met Thr 610 615 620

Trp Met Glu Trp Asp Arg Glu Ile Asn Asn Tyr Thr Ser Ile Ile Tyr 625 630 635 640

Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu 645 650 655

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Glu Arg Ala Leu Leu 850